| Project Title   | Funding     | Strategic Plan Objective | Institution                             |  |
|---|-------------|--------------------------|---|--|
| Senetic basis of autism   | \$4,000,000 | Q3.L.B                   | Cold Spring Harbor Laboratory           |  |
| Autism Genetics, Phase II: Increasing Representation of Human Diversity         | \$2,728,166 | Q3.S.D                   | University of California, Los Angeles   |  |
| Genetic Epidemiology of Complex Traits  | \$808,519   | Q3.L.B                   | National Institutes of Health           |  |
| Autism genetics: homozygosity mapping and functional alidation                  | \$765,736   | Q3.L.B                   | CHILDREN'S HOSPITAL CORPORATION         |  |
| /4-The Autism Sequencing Consortium: Autism gene iscovery in >20,000 exomes     | \$720,372   | Q3.S.A                   | ICAHN SCHOOL OF MEDICINE AT MOUNT SINAI |  |
| /4 The Autism Sequencing Consortium: Autism gene liscovery in >20,000 exomes    | \$674,849   | Q3.S.A                   | UNIVERSITY OF CALIFORNIA, SAN FRANCISCO |  |
| Population-Based Autism Genetics & Environment Study                            | \$655,813   | Q3.L.D                   | ICAHN SCHOOL OF MEDICINE AT MOUNT SINAI |  |
| poradic Mutations and Autism Spectrum Disorders                                 | \$647,900   | Q3.S.A                   | University of Washington                |  |
| tapid Phenotyping for Rare Variant Discovery in Autism                          | \$589,746   | Q3.S.A                   | University of California, Los Angeles   |  |
| accelerating Autism Genetics via Whole Population accertainment in Denmark      | \$504,821   | Q3.L.B                   | Broad Institute, Inc.                   |  |
| /4-The Autism Sequencing Consortium: Autism gene iscovery in >20,000 exomes     | \$415,893   | Q3.S.A                   | BROAD INSTITUTE, INC.                   |  |
| lieces of the Puzzle: Uncovering the Genetics of Autism                         | \$374,636   | Q3.L.B                   | Broad Institute, Inc.                   |  |
| utism subtypes by gene characterization   | \$368,757   | Q3.S.A                   | University of Washington                |  |
| lequencing Female-enriched Multiplex Autism Families FEMFs)                     | \$353,610   | Q3.L.B                   | Johns Hopkins University                |  |
| lovel Statistical methods for DNA Sequencing Data, nd applications to Autism.   | \$318,575   | Q3.L.B                   | Columbia University                     |  |
| /3 - Sequencing Autism Spectrum Disorder Extended Pedigrees                     | \$298,000   | Q3.L.B                   | University of Utah                      |  |
| Senetic basis of phenotypic variability in 16p11.2 eletion or duplication       | \$281,679   | Q3.L.B                   | University of Washington                |  |
| flutations in noncoding DNA and the missing heritability f autism               | \$275,697   | Q3.L.B                   | University of California, San Diego     |  |
| 0/4 - The Autism Sequencing Consortium: Autism gene liscovery in >20,000 exomes | \$263,975   | Q3.S.A                   | University of Pittsburgh                |  |
| Complex Genetic Architecture of Chromosomal<br>berrations in Autism             | \$248,999   | Q3.L.B                   | Massachusetts General Hospital          |  |
| /3 Sequencing Autism Spectrum Disorder Extended ledigrees                       | \$231,750   | Q3.L.B                   | University of Washington                |  |
| lon-Coding RNAs in Autism   | \$205,365   | Q3.Other                 | UNIVERSITY OF SOUTHERN CALIFORNIA       |  |
| itochondria and the etiology of autism  | \$175,000   | Q3.L.B                   | Children's Hospital of Philadelphia     |  |
| denomic influences on development and outcomes in a string for autism           | \$171,665   | Q3.L.B                   | University of Alberta                   |  |
| /3-Sequencing Autism Spectrum Disorder Extended ledigrees                       | \$160,000   | Q3.L.B                   | UNIVERSITY OF PENNSYLVANIA              |  |

| Project Title  | Funding   | Strategic Plan Objective | Institution                                    |  |
|--|-----------|--------------------------|--|--|
| Genomic influences on developmental course and<br>outcome in Infants at risk of ASD: A Baby Siblings<br>Research Consortium (BSRC) Study | \$148,802 | Q3.S.A                   | University of Alberta                          |  |
| Investigating the role of somatic mutations in autism spectrum disorders   | \$139,366 | Q3.L.B                   | Oregon Health & Science University             |  |
| Whole-exome sequencing to identify causative genes for autism  | \$134,203 | Q3.L.B                   | Rockefeller University                         |  |
| Autism Spectrum Disorders: Genomes to Outcomes   | \$133,500 | Q3.L.B                   | The Hospital for Sick Children                 |  |
| dentification and analysis of functional networks<br>perturbed in autism   | \$125,000 | Q3.L.B                   | Columbia University                            |  |
| Genome-wide analysis of cis-regulatory elements in autism  | \$125,000 | Q3.L.B                   | Washington University in St. Louis             |  |
| Computational tools to analyze SNP data from patients with mental illness  | \$120,877 | Q3.L.B                   | PARTEK, INC.                                   |  |
| Elucidating pathogenic mutations disrupting RNA regulation in autism   | \$112,500 | Q3.L.B                   | Columbia University                            |  |
| Genome Sequencing pilot of Simons Simplex Collection   | \$100,643 | Q3.L.B                   | University of Washington                       |  |
| Cryptic chromosomal aberrations contributing to autism   | \$65,125  | Q3.L.B                   | Massachusetts General Hospital                 |  |
| Dosage effects of DUF1220 gene subtype CON1 in autism  | \$62,500  | Q3.L.B                   | University of Colorado, Denver                 |  |
| Dosage effects of 22q11 region on autism-relevant neural systems   | \$60,000  | Q3.S.A                   | University of California, Los Angeles          |  |
| Discovery and Functional Characterization of Gene Regulatory Networks (GRNs) of Autism Risk Genes  | \$57,900  | Q3.Other                 | Children's Hospital Los Angeles                |  |
| Genomic profiling of autism families using whole-<br>genome sequencing   | \$45,280  | Q3.L.B                   | Institut Pasteur                               |  |
| An exploration of genetic and behavioral variables in<br>Autism Spectrum Disorder  | \$45,000  | Q3.S.A                   | Center for Autism and Related Disorders (CARD) |  |
| The genetic basis underlying the phenotype neterogeneity of the 16p11.2 CNV  | \$37,550  | Q3.S.A                   | University of Washington                       |  |
| Sequence-based discovery of genes with pleiotropic effects across diagnostic boundaries and throughout the ifespan                       | \$29,995  | Q3.L.B                   | Massachusetts General Hospital                 |  |
| Role of the Intestinal Microbiome in Children with Autism  | \$29,000  | Q3.S.I                   | Massachusetts General Hospital                 |  |
| Simons Simplex Collection support grant  | \$13,200  | Q3.L.B                   | University of California, Los Angeles          |  |
| Simons Simplex Collection support grant  | \$10,000  | Q3.L.B                   | University of Washington                       |  |
| imons Simplex Collection support grant   | \$10,000  | Q3.L.B                   | Vanderbilt University                          |  |
| Simons Simplex Collection support grant  | \$10,000  | Q3.L.B                   | Baylor College of Medicine                     |  |
| Simons Simplex Collection support grant  | \$10,000  | Q3.L.B                   | University of Missouri                         |  |

| Project Title   | Funding  | Strategic Plan Objective | Institution   |  |
|---|----------|--------------------------|---|--|
| Simons Simplex Collection support grant   | \$10,000 | Q3.L.B                   | McGill University Health Centre- Montreal Children's Hospital |  |
| Simons Simplex Collection support grant   | \$10,000 | Q3.L.B                   | Yale University   |  |
| Simons Simplex Collection support grant   | \$4,134  | Q3.L.B                   | University of Illinois at Chicago                             |  |
| Whole exome sequencing of Simons Simplex Collection quads   | \$0      | Q3.L.B                   | University of Washington                                      |  |
| A history of behavioral genetics  | \$0      | Q3.Other                 | University of Pittsburgh                                      |  |
| Simons Simplex Collection support grant   | \$0      | Q3.L.B                   | Boston Children's Hospital                                    |  |
| Genomic influences on development and outcomes in Infants at risk of ASD                          | \$0      | Q3.S.A                   | University of Alberta   |  |
| A genome-wide search for autism genes in the SSC UCLA   | \$0      | Q3.L.B                   | University of California, Los Angeles                         |  |
| Simons Simplex Collection support grant   | \$0      | Q3.L.B                   | Weill Cornell Medical College                                 |  |
| Simons Simplex Collection support grant   | \$0      | Q3.L.B                   | Columbia University   |  |
| Simons Simplex Collection support grant   | \$0      | Q3.L.B                   | Emory University  |  |
| Identifying genetic variants on the Y chromosome of males with autism                             | \$0      | Q3.L.B                   | The Hospital for Sick Children                                |  |
| Identification of functional networks perturbed in autism   | \$0      | Q3.L.B                   | Columbia University   |  |
| Illumina, Inc.  | \$0      | Q3.L.B                   | Illumina, Inc.  |  |
| High-throughput Screening of Novel Trinucleotide<br>Repeat Expansion in Autism Spectrum Disorders | \$0      | Q3.L.B                   | The Hospital for Sick Children                                |  |
| Identifying Patterns of Genetic Variants Conferring Risk for Neurodevelopmental Disorders         | \$0      | Q3.L.B                   | Pennsylvania State University                                 |  |